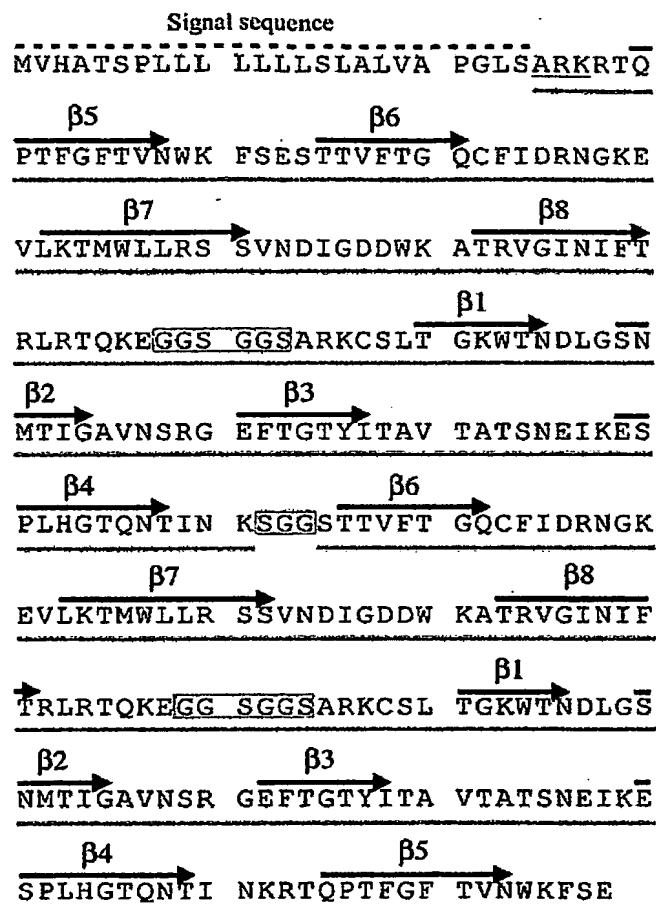


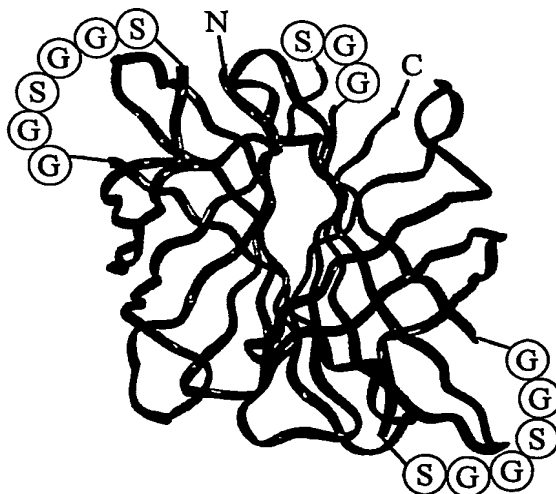
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Fig. 1

A

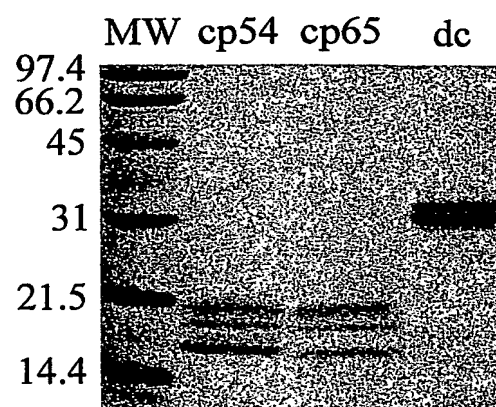


B



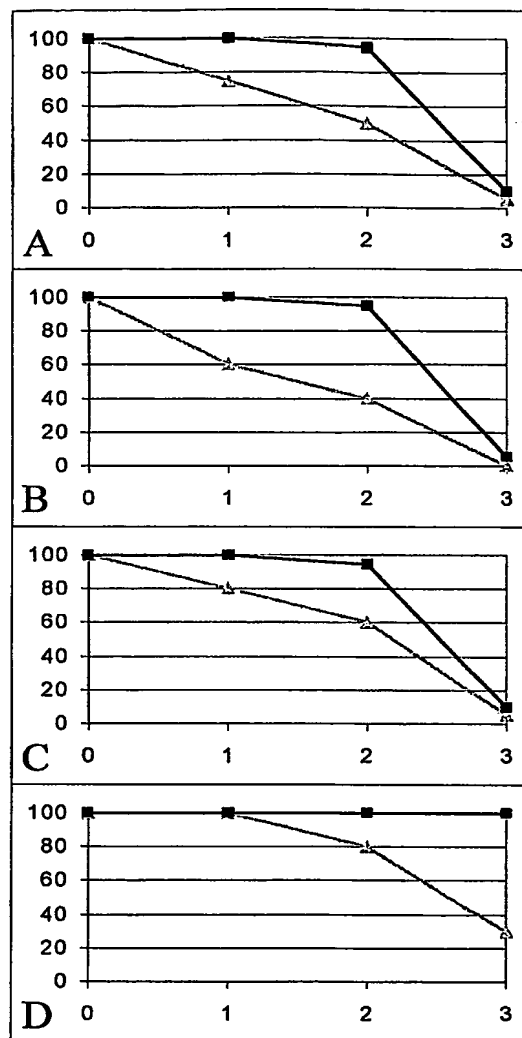
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Fig. 2



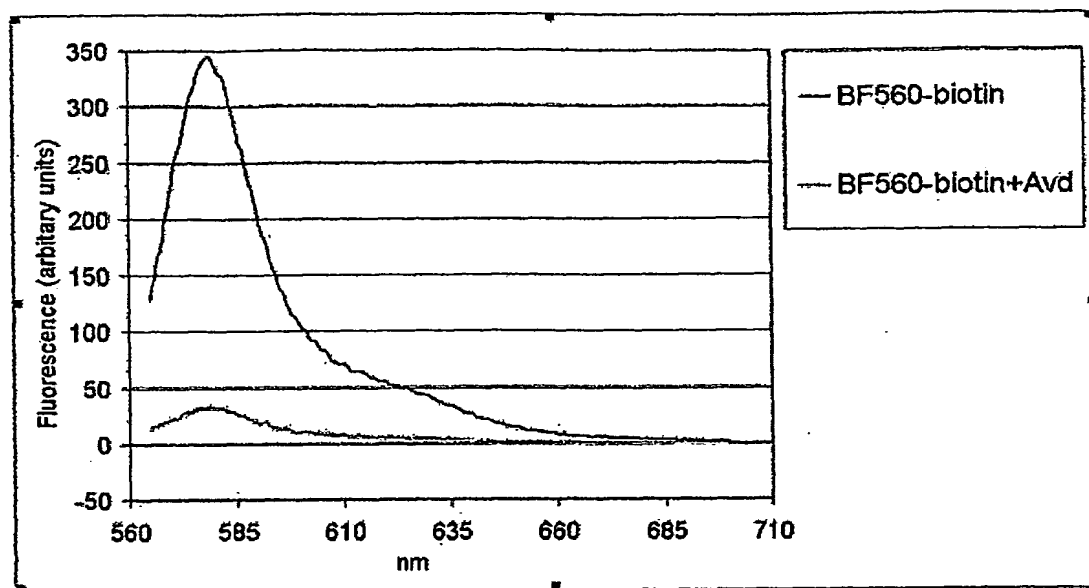
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Fig. 3



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Fig. 4



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Fig. 5

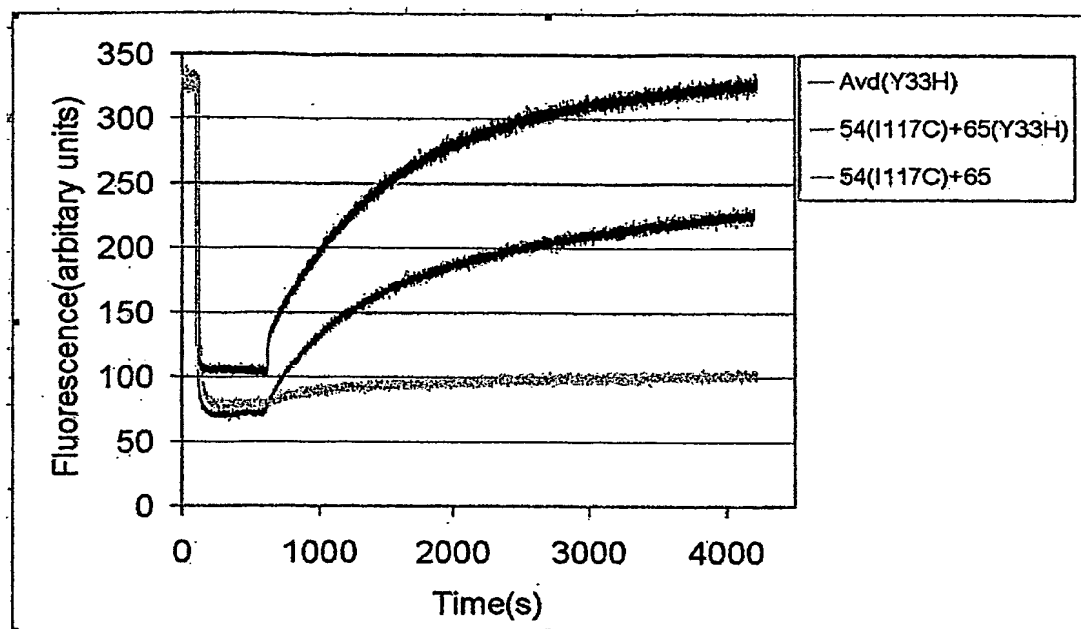


Fig. 6

ATGGTGCACGCAACCTCCCCGCTGCTGCTGCTGCTGCT
GCTCAGCCTGGCTCTGGTGGCTCCCGGCCTCTCTGCCA
GGAAGAGGACCCAGCCCACCTTTGGCTTCACCGTCAAT
TGGAAGTTTTTCAGAGTCCACCACTGTCTTCACGGGCCA
GTGCTTCATAGACAGGAATGGGAAGGAGGTCCTGAAG
ACCATGTGGCTGCTGCGGTCAAGTGTTAATGACATTGG
TGATGACTGGAAAGCTACCAGGGTCGGCATCAACATC
TTCCTCGCCTGCGCACACAGAAGGAGGGAGGCTCCG
GAGGCTCCGCCAGAAAGTGCTCGCTGACTGGGAAATG
GACCAACGATCTGGGCTCCAACATGACCATCGGGGCT
GTGAACAGCAGAGGTGAATTCACAGGCACCTACATCA
CAGCCGTAACAGCCACATCAAATGAGATCAAAGAGTC
ACCACTGCATGGGACACAAAACACCATCAACAAGTCC
GGCGGATCCACCACTGTCTTCACGGGCCAGTGCTTCAT
AGACAGGAATGGGAAGGAGGTCCTGAAGACCATGTGG
CTGCTGCGGTCAAGTGTTAATGACATTGGTGATGACTG
GAAAGCTACCAGGGTCGGCATCAACATCTTCACTCGCC
TGCGCACACAGAAGGAGGGAGGCTCCGGAGGCTCCGC
CAGAAAGTGCTCGCTGACTGGGAAATGGACCAACGAT
CTGGGCTCCAACATGACCATCGGGGCTGTGAACAGCA
GAGGTGAATTCACAGGCACCTACATCACAGCCGTAAC
AGCCACATCAAATGAGATCAAAGAGTCACCACTGCAT
GGGACACAAAACACCATCAACAAGAGGACCCAGCCCA
CCTTTGGCTTCACCGTCAATTGGAAGTTTTTCAGAGGGA
GGTTCGGGATCGGGATCCGGCTCTGGCAGCGGCAGGA
CCCAGCCCACCTTTGGCTTCACCGTCAATTGGAAGTTT
TCAGAGTCCACCACTGTCTTCACGGGCCAGTGCTTCAT
AGACAGGAATGGGAAGGAGGTCCTGAAGACCATGTGG
CTGCTGCGGTCAAGTGTTAATGACATTGGTGATGACTG
GAAAGCTACCAGGGTCGGCATCAACATCTTCACTCGCC
TGCGCACACAGAAGGAGGGAGGCTCCGGAGGCTCCGC
CAGAAAGTGCTCGCTGACTGGGAAATGGACCAACGAT
CTGGGCTCCAACATGACCATCGGGGCTGTGAACAGCA
GAGGTGAATTCACAGGCACCTACATCACAGCCGTAAC
AGCCACATCAAATGAGATCAAAGAGTCACCACTGCAT
GGGACACAAAACACCATCAACAAGTCCGGCGGATCCA
CCTGTCTTCACGGGCCAGTGCTTCATAGACAGGAAT
GGGAAGGAGGTCCTGAAGACCATGTGGCTGCTGCGGT
CAAGTGTTAATGACATTGGTGATGACTGGAAAGCTAC
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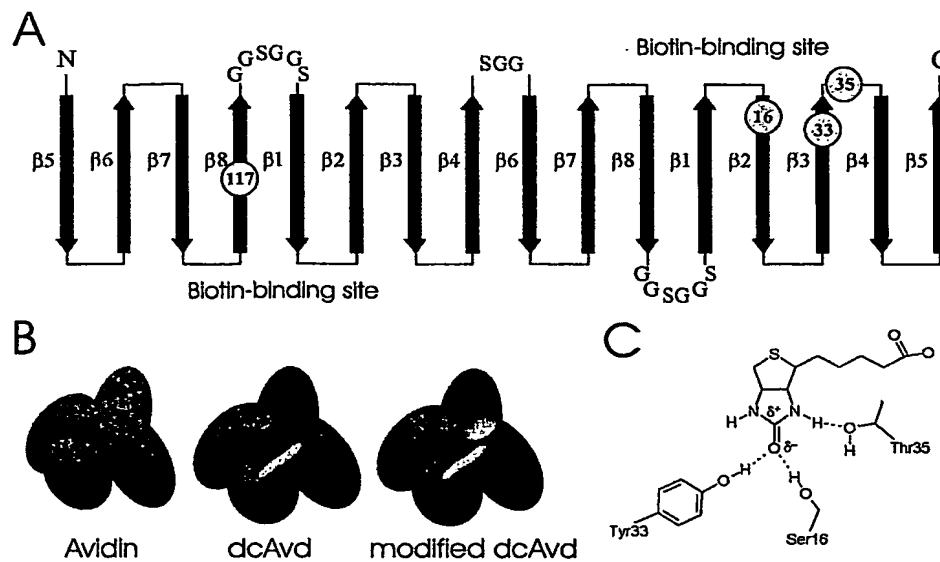
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Fig. 7

ATGGTGCACGCAACCTCCCCGCTGCTGCTGCTGCTGCTGCTCA
GCCTGGCTCTGGTGGCTCCCGGCTCTCTGCCAGGAAGAGGAC
CCAGCCCACCTTTTGGCTTCACCGTCAATTGGAAGTTTTTCAGAG
TCCACCCTGTCTTCACGGGCCAGTGCTTCATAGACAGGAATG
GGAAGGAGGTCCTGAAGACCATGTGGCTGCTGCGGTCAAGTGT
TAATGACATTGGTGATGACTGGAAAGCTACCAGGGTCGGCATC
AACATCTTCACTCGCCTGCGCACACAGAAGGAGGGAGGCTCCG
GAGGCTCCGCCAGAAAGTGCTCGCTGACTGGGAAATGGACCAA
CGATCTGGGCTCCAACATGACCATCGGGGCTGTGAACAGCAGA
GGTGAATTCACAGGCACCTACATCACAGCCGTAACAGCCACAT
CAAATGAGATCAAAGAGTCACCACTGCATGGGACACAAAACAC
CATCAACAAGTCCGGCGGATCCACCACTGTCTTCACGGGCCAG
TGCTTCATAGACAGGAATGGGAAGGAGGTCCTGAAGACCATGT
GGCTGCTGCGGTCAAGTGTTAATGACATTGGTGATGACTGGAA
AGCTACCAGGGTCGGCATCAACATCTTCACTCGCCTGCGCACA
CAGAAGGAGGGAGGCTCCGGAGGCTCCGCCAGAAAGTGCTCGC
TGACTGGGAAATGGACCAACGATCTGGGCTCCAACATGACCAT
CGGGGCTGTGAACAGCAGAGGTGAATTCACAGGCACCTACATC
ACAGCCGTAACAGCCACATCAAATGAGATCAAAGAGTCACCAC
TGCATGGGACACAAAACACCATCAACAAGAGGACCCAGCCCAC
CTTTGGCTTCACCGTCAATTGGAAGTTTTTCAGAGTGA

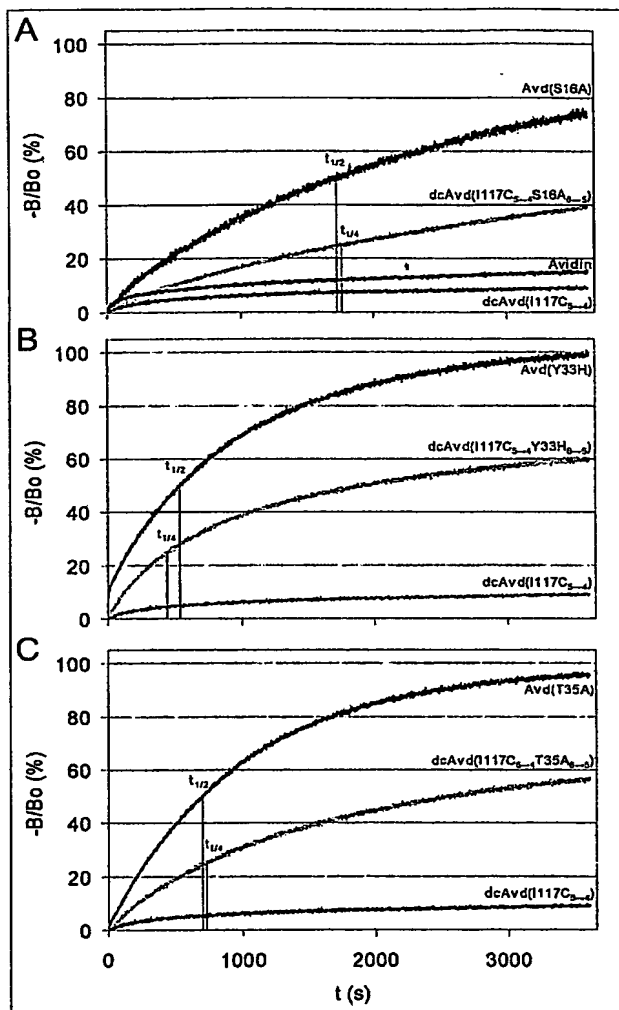
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Fig. 8



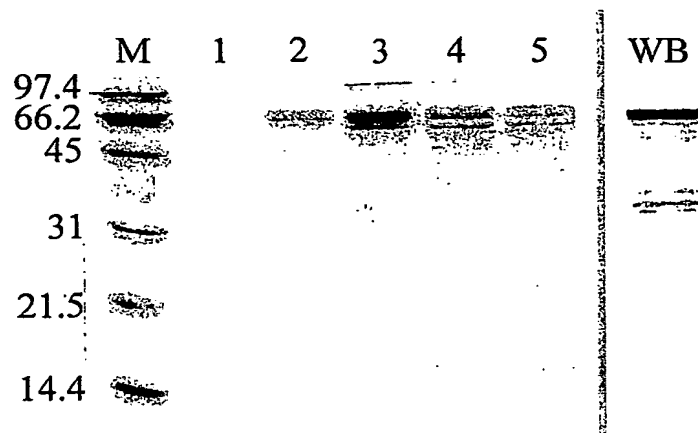
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Fig. 9



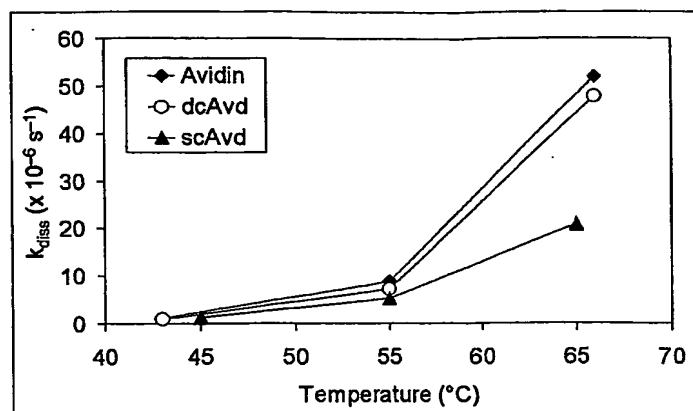
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FIG. 10



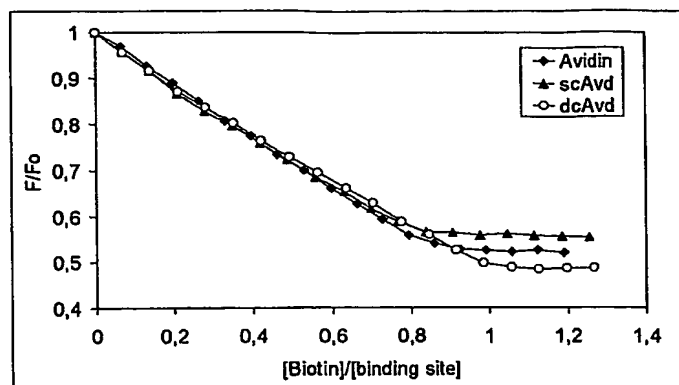
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FIG. 11



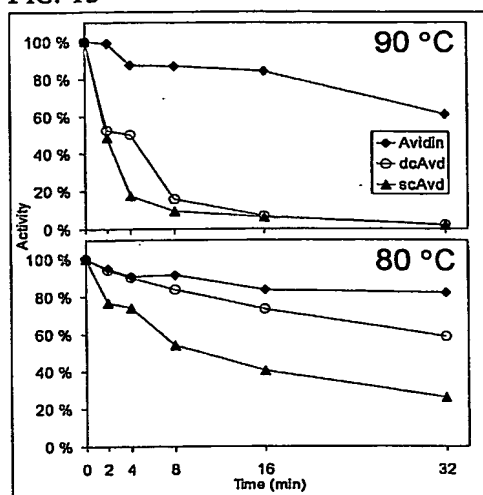
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FIG. 12



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FIG. 13



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Fig. 14

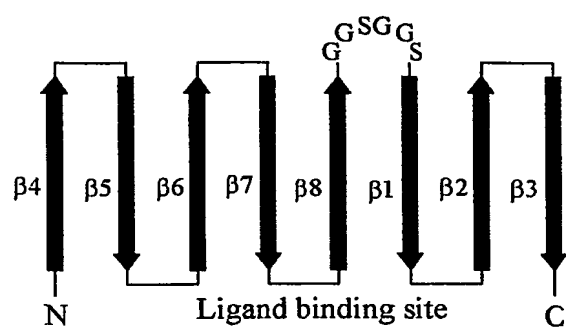
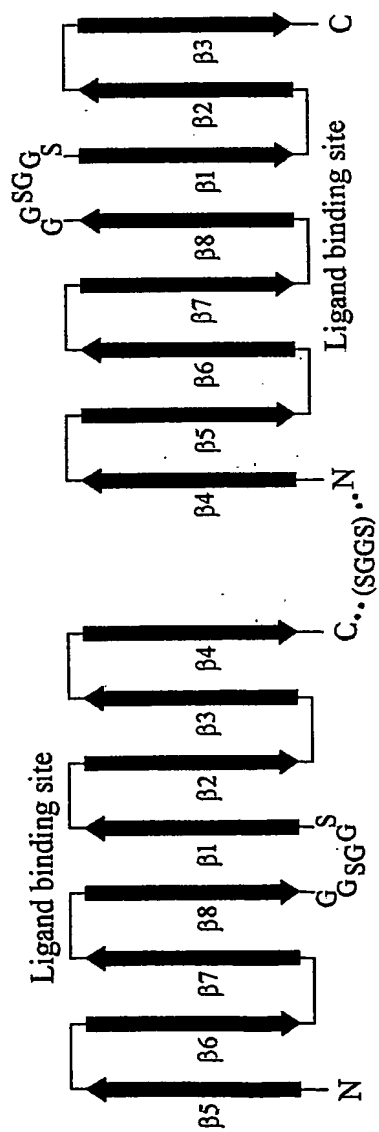
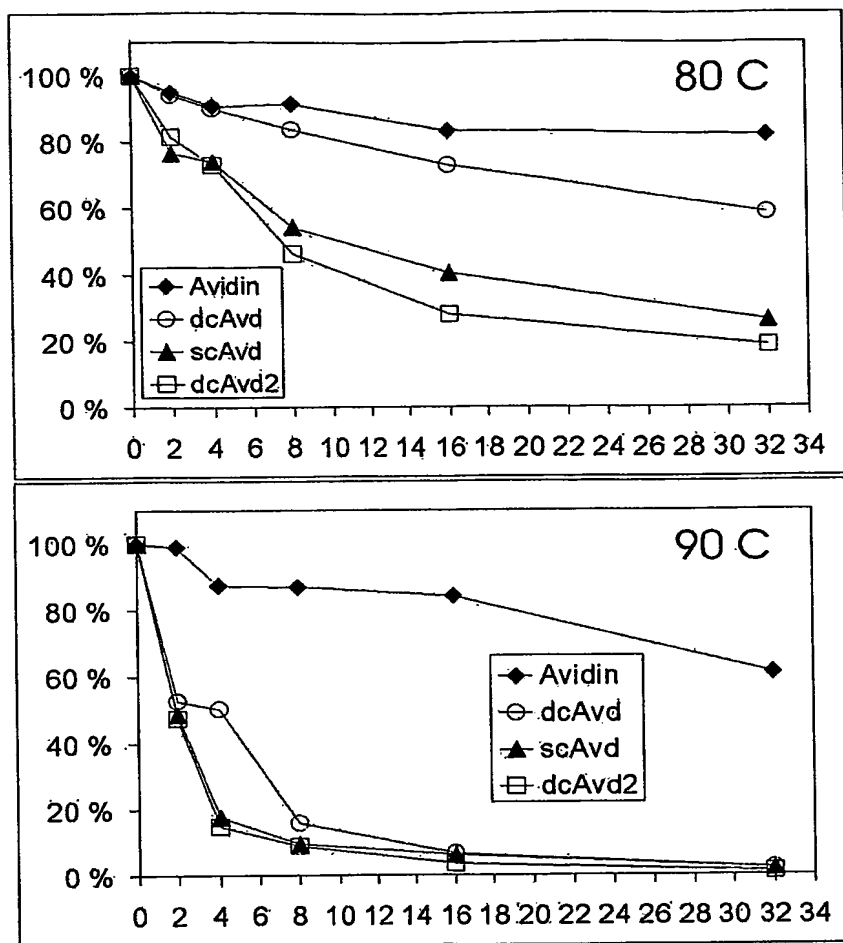


Fig 15.



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Fig. 16



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Fig. 17

1 MNKPSKFALP LAFAAVTASG VASAGTQPTF GFTVNWKFSE
STTVFTGQCF IDRNGKEVLK
61 TMWLLRSSVN DIGDDWKATR VGINIFTRLR TQKEGGSGGS
ARKCSLTGKW TNDLGSNMTI
121 GAVNSRGEFT GTYITAVTAT SNEIKESPLH GTQNTINKSG
GSKESPLHGT QNTINKRTQP
181 TFGFTVNWKF SESTTVFTGQ CFIDRNGKEV LKTMWLLRSS
VNDIGDDWKA TRVGINIFTR
241 LRTQKEGGSG GSARKCSLTG KWTNDLGSNM TIGAVNSRGE
FTGTYITAVT

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Fig. 18

```
1      ATGAACAAAC CCTCCAAATT CGCTCTGCCG CTTGCCTTCG
CGGCCGTTAC GGCCTCTGGT
61      GTTGCCTCGG CTGGTACCCA GCCCACCTTT GGCTTCACCG
TCAATTGGAA GTTTTCAGAG
121     TCCACCACTG TCTTCACGGG CCAGTGCTTC ATAGACAGGA
ATGGGAAGGA GGTCCCTGAAG
181     ACCATGTGGC TGCTGCGGTC AAGTGTTAAT GACATTGGTG
ATGACTGGAA AGCTACCAGG
241     GTCGGCATCA ACATCTTCAC TCGCCTGCGC ACACAGAAGG
AGGGAGGCTC CGGAGGCTCC
301     GCCAGAAAGT GCTCGCTGAC TGGGAAATGG ACCAACGATC
TGGGCTCCAA CATGACCATC
361     GGGGCTGTGA ACAGCAGAGG TGAATTCACA GGCACCTACA
TCACAGCCGT AACAGCCACA
421     TCAAATGAGA TCAAAGAGTC ACCACTGCAT GGGACACAAA
ACACCATCAA CAAGTCCGGC
481     GGATCCAAAG AGTCACCACT GCATGGGACA CAAAACACCA
TCAACAAGAG GACCCAGCCC
541     ACCTTTGGCT TCACCGTCAA TTGGAAGTTT TCAGAGTCCA
CCTACTGTCTT CACGGGCCAG
601     TGCTTCATAG ACAGGAATGG GAAGGAGGTC CTGAAGACCA
TGTGGCTGCT GCGGTCAAGT
661     GTTAATGACA TTGGTGATGA CTGGAAAGCT ACCAGGGTCG
GCATCAACAT CTTCCTCGC
721     CTGCGCACAC AGAAGGAGGG AGGCTCCGGA GGCTCCGCCA
GAAAGTGCTC GCTGACTGGG
781     AAATGGACCA ACATCTGGG CTCCAACATG ACCATCGGGG
CTGTGAACAG CAGAGGTGAA
841     TTCACAGGCA CCTACATCAC AGCCGTAACA TAA
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